

Mailbox:

12 E12

Room: 12D07

SEARCH REQUEST FORM

U.S. DEPARTMENT OF COMMERCE
Patent and Trademark Office

82244 *

Requestor's
Name:

Anne-Marie Falk

Serial

Number:

09/435,471

Date:

12/12/02

Phone:

306-9155

Art Unit:

1632

Search Topic:

Please write a detailed statement of search topic. Describe specifically as possible the subject matter to be searched. Define any terms that may have a special meaning. Give examples or relevant citations, authors, keywords, etc., if known. For sequences, please attach a copy of the sequence. You may include a copy of the broadest and/or most relevant claim(s).

- Please search SEQ ID No. 9.

- Search: commercial databases and
issued database.
(and pre-grant pubs database if
there is such a thing)

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Point of Contact
P. Sheppard
Telephone number: (703) 305-4489

NA 9-33

STAFF USE ONLY

Date completed:

Searcher:

Terminal time:

Elapsed time:

CPU time:

Total time:

Number of Searches:

Number of Databases:

Search Site

STIC

CM-1

Pre-S

Type of Search

N.A. Sequence

A.A. Sequence

Structure

Bibliographic

Vendors

IG

STN

Dialog

APS

Geninfo

SDC

DARC/Questel

Other

WEST Search History

DATE: Thursday, December 12, 2002

Set Name Query
side by side

Hit Count Set Name
result set

DB=USPT; PLUR=YES; OP=ADJ

L3 instability element and (PKC or protein kinase)

2 L3

L2 mRNA and transcription

18481 L2

L1 mRNA and instability element

16 L1

END OF SEARCH HISTORY

*Amz
12/12/02*

?ds

Set	Items	Description
S1	213	INSTABILITY(W) ELEMENT
S2	187	S1 AND MRNA
S3	63	RD (unique items)
S4	3	S3 AND GLUCOSE
S5	29799	MRNA AND ELEMENT AND TRANSCRIPTION
S6	1117	S5 AND GLUCOSE
S7	195	S6 AND (STABILITY OR INSTABILITY)
S8	149	RD (unique items)
S9	108	S8 AND (INCREASE OR DECREASE)
S10	0	S9 AND (MRNA NEAR STABILITY)
S11	0	S9 AND MRNA (NEAR) STABILITY
S12	41	S9 AND MRNA (W) STABILITY
?		

Amr
12/12/02

4/3,AB/1 (Item 1 from file: 5)
DIALOG(R) File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

13536523 BIOSIS NO.: 200200165344

**Insulin-regulated inclusion of a protein kinase C betaII-specific exon
introduces an instability element responding to high glucose
concentrations in vascular smooth muscle cells.**

AUTHOR: Cooper Denise R(a); Patel Niketa A; Chappell David S; Mebert Konrad
; Mancu Daniel; Miller Laura A; Duong Tram Thuy; Sridhar Rajagopalan;
Watson James E; Eichler Duane C

AUTHOR ADDRESS: (a)Research Service, J.A. Haley Veterans Hospital, 13000
Bruce B. Downs Blvd., Tampa, FL, 33612**USA

JOURNAL: Molecular Biology of the Cell 12 (Supplement):p229a Nov, 2001

MEDIUM: print

CONFERENCE/MEETING: 41st Annual Meeting of the American Society for Cell
Biology Washington DC, USA December 08-12, 2001

ISSN: 1059-1524

RECORD TYPE: Citation

LANGUAGE: English

2001

Am2

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RESULT 8
US-08-388-672A-5
; Sequence 5, Application US/08388672A
; Patent No. 5795961
; GENERAL INFORMATION:
;   APPLICANT: Wallace, T. Paul
;   APPLICANT: Harris, William J.
;   APPLICANT: Carr, Frank J.
;   APPLICANT: Old, Lloyd J.
;   APPLICANT: Welt, Sydney
;   APPLICANT: Kitamura, Kunio
;   TITLE OF INVENTION: Recombinant Human Anti-Lewis B
;   TITLE OF INVENTION: Antibodies
;   NUMBER OF SEQUENCES: 25
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: Felte and Lynch
;     STREET: 805 Third Avenue
;     CITY: New York
;     STATE: New York
;     COUNTRY: U.S.A.
;     ZIP: 10022
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: Floppy disk
;     COMPUTER: IBM PC compatible
;     OPERATING SYSTEM: PC-DOS/MS-DOS
;     SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/08/388,672A
;     FILING DATE: 14-FEB-1995
;     CLASSIFICATION:
;   ATTORNEY/AGENT INFORMATION:
;     NAME: Hanson, No. 5795961man D.
;     REGISTRATION NUMBER: 30,946
;     REFERENCE/DOCKET NUMBER: LUD 5409
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: 212-688-9200
;     TELEFAX: 212-838-3884
;   INFORMATION FOR SEQ ID NO: 5:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH: 37 base pairs
;
;     TYPE: nucleic acid
;     STRANDEDNESS: unknown
;     TOPOLOGY: unknown
;     MOLECULE TYPE: DNA (genomic)
US-08-388-672A-5

Query Match      53.3%; Score 17.6; DB 1; Length 37;
Best Local Similarity 65.4%; Pred. No. 13;
Matches 17; Conservative 5; Mismatches 4; Indels 0; Gaps 0;

Qy      6 CTAGTCGACAAGAGTTTGTCTAGTGGG 31
        ||||| ||| |:: |::|
Db      2 CTAGTCGACATGAGGTGYYTGYSG 27
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